

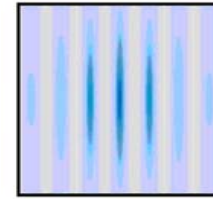
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The Atomic Pair Distribution Function Method

Getting to know your atomic
neighborhood

Introduction

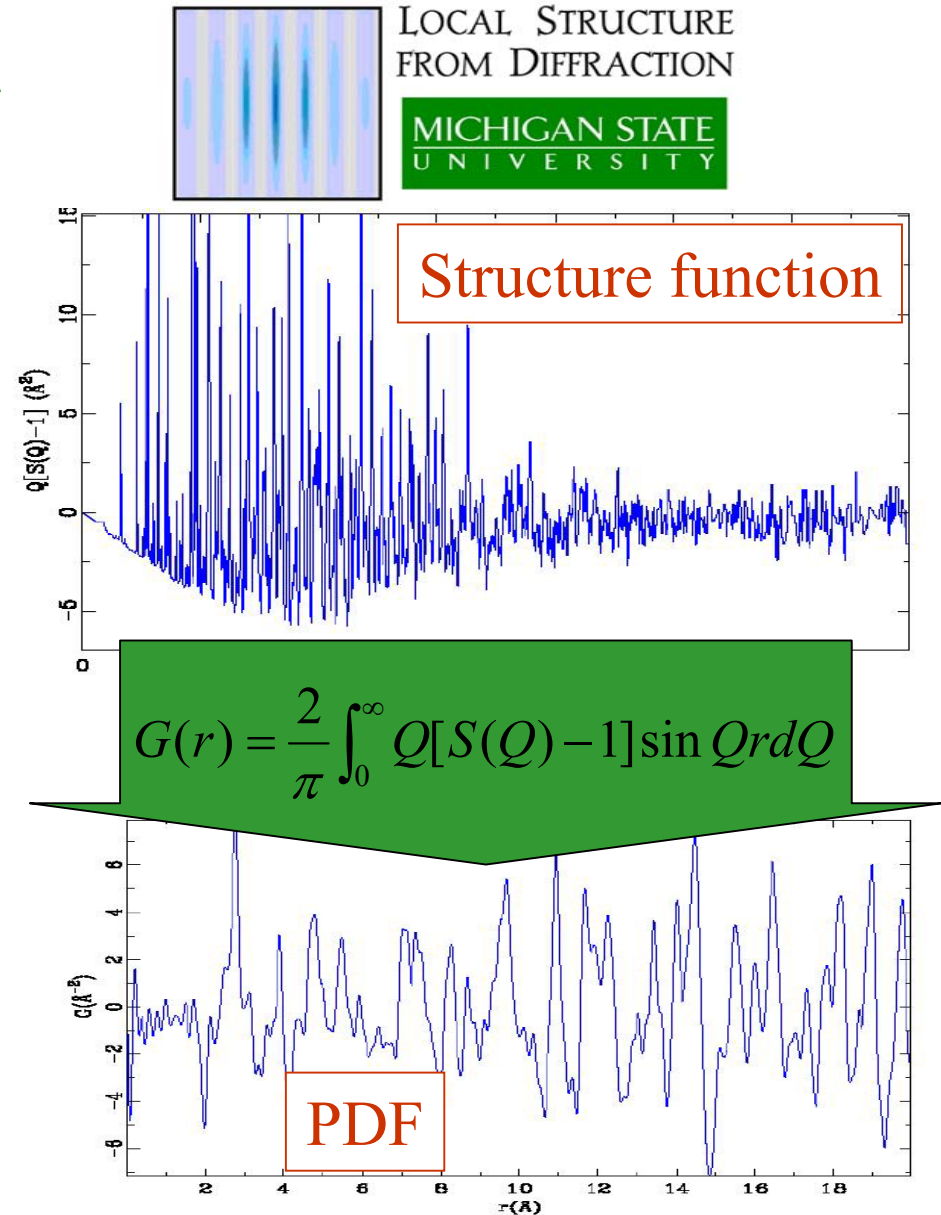
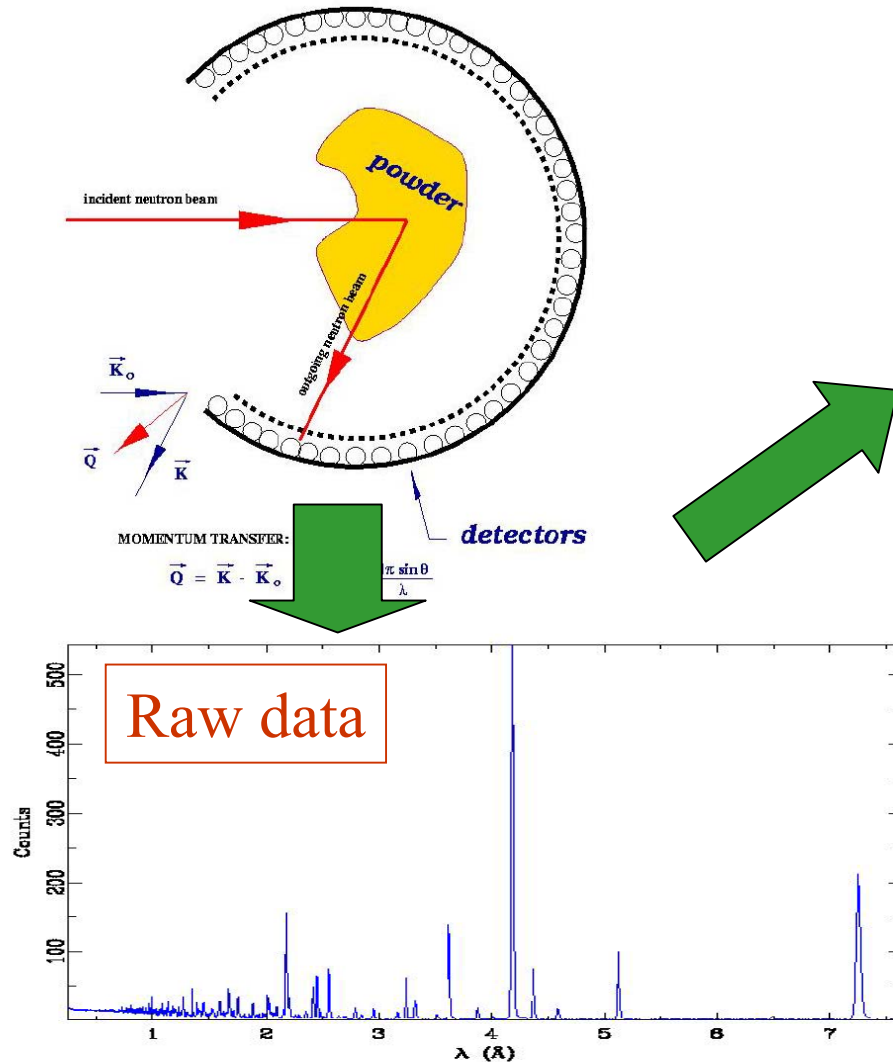


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- Modern materials are often disordered.
- Standard crystallographic methods lose the aperiodic (disorder) information.
- We would like to be able to **sit on an atom and look at our neighborhood**.
- The **PDF method** allows us to do that (see next slide):
 - First we do a neutron or x-ray diffraction experiment
 - Then we correct the data for experimental effects
 - Then we Fourier transform the data to real-space

Obtaining the PDF

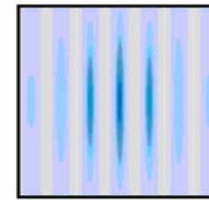
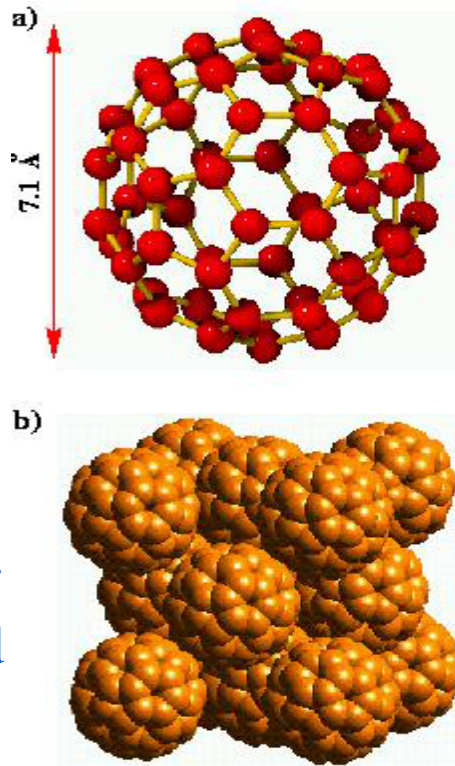


What is the PDF?

- (a) The red ball is a C_{60} molecule. C_{60} forms a solid by the molecules clustering (b). The scattering and PDF are shown in (c) and (d) respectively.

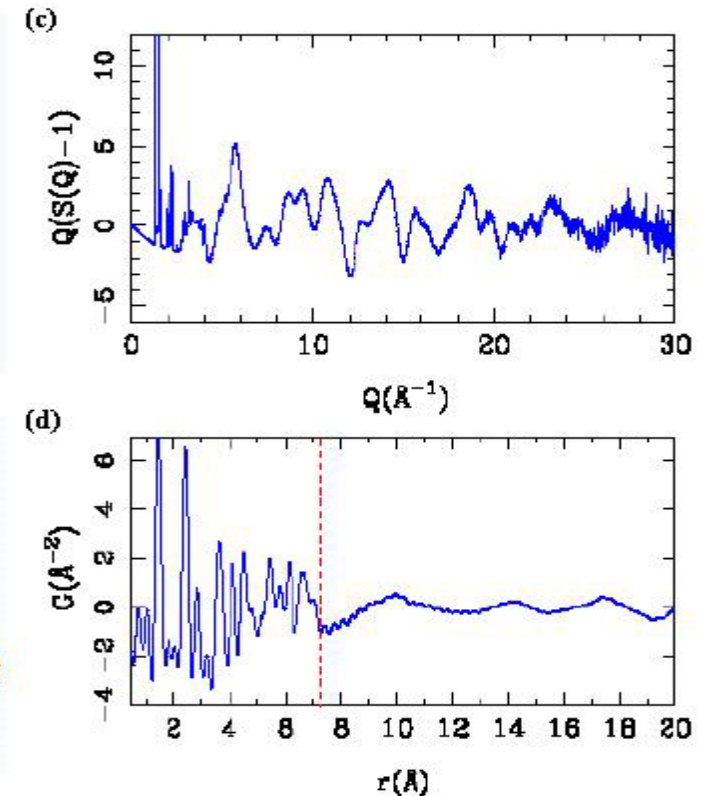
- Sit on an atom and look at your neighborhood. The nearest neighbor is at 1.4Å distance, the second neighbor at 2.2Å and so on. There are sharp peaks in $G(r)$ at these positions. This is the structural information in the PDF.

- There are no sharp peaks beyond 7.1Å, the diameter of the ball because the balls are spinning with respect to each other. The PDF can see this.

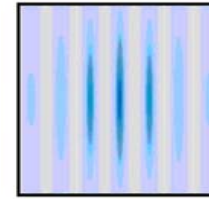


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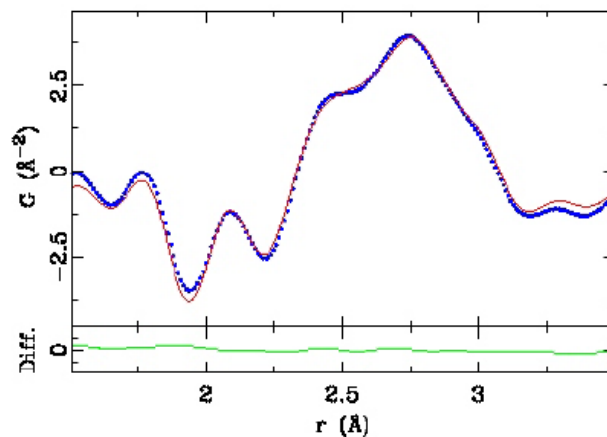
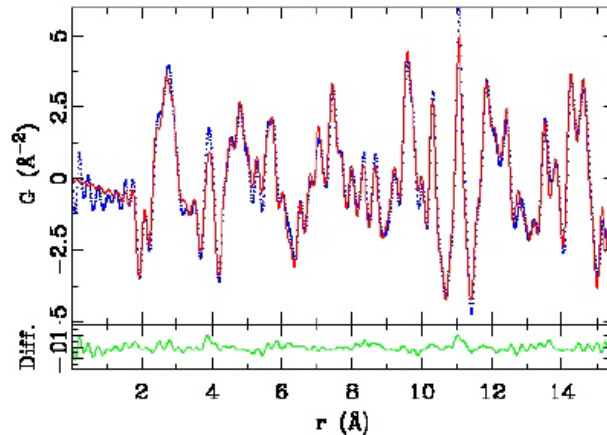


Full Profile PDF fitting using PDFFIT



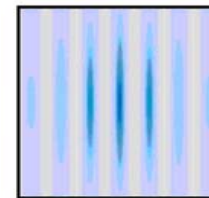
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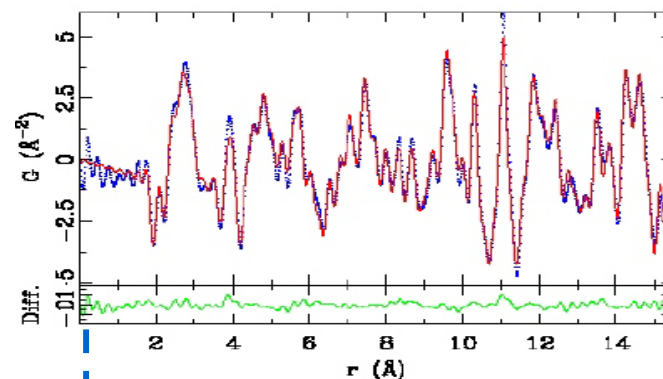
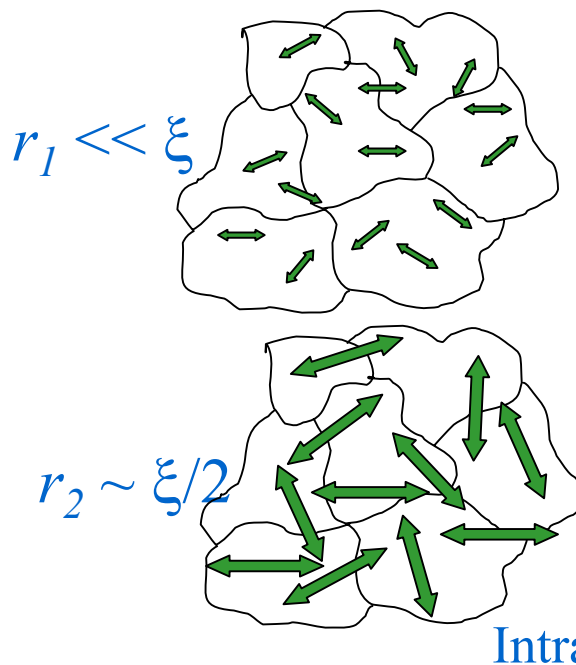
- We extract information from the PDF by **fitting structural models to the data**
- We use Full-profile least-squares fitting of the PDF using the program PDFfit
- The **red** line is the PDF from the model, the **blue** line the data, the **green** line the difference.
- The data are neutron data from LaMnO_3 collected 10K at IPNS, Argonne National Laboratory, IL.
- **Ref: Proffen et al., Phys. Rev. B 60, 9973 (1999).**

Observing Domains in the PDF



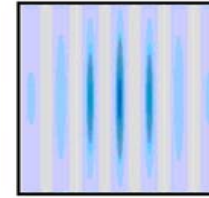
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The PDF gives different information on different length-scales. We can see the structure **within a domain** at low- r and **between domains** at high- r .

Goodbye



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- Thanks for the visit!